RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/550,788_

Source: _PU/10_

Date Processed by STIC: _10/7/05_

ENTERED

Raw Sequence Listing before editing
(for reference only)

PCT

RAW SEQUENCE LISTING   DATE: 10/07/2005
PATENT APPLICATION: **US/10/550,788**   TIME: 09:31:06

Input Set : **A:\Sequence Listing 2005-1542A.txt**
Output Set: **N:\CRF4\10072005\J550788.raw**

```
   3 <110> APPLICANT: President of National Rehabilitation Center for Persons with
Disabilities
   4        Hitachi Instruments Service Co., Lt.
   5        KATO, Seishi            Method
   7 <120> TITLE OF INVENTION:(Mehod) for synthesizing cDNA
   9 <130> FILE REFERENCE: 04-F-018PCT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/550,788
C--> 12 <141> CURRENT FILING DATE: 2005-09-28
  14 <150> PRIOR APPLICATION NUMBER: JP2003-91373
  15 <151> PRIOR FILING DATE: 2003-03-28
  17 <160> NUMBER OF SEQ ID NOS: 2
  19 <170> SOFTWARE: PatentIn version 3.1
```

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

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  92 <210> SEQ ID NO: 2
  93 <211> LENGTH: 3357
  94 <212> TYPE: DNA
  95 <213> ORGANISM: Artificial
  97 <220> FEATURE:
  98 <223> OTHER INFORMATION: pGCAP10: Modified from pGCAP1 of SEQ ID No. 1
 100 <220> FEATURE:
 101 <221> NAME/KEY: misc_feature
 102 <223> OTHER INFORMATION: Circular polynucleotide
 104 <400> SEQUENCE: 2
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 107 gctccccagc aggcagaagt atgcaaagca tgcatctcaa ttagtcagca accatagtcc     180
 108 cgcccctaac tccgcccatc ccgcccctaa ctccgcccag ttccgcccat tctccgcccc     240
 109 atggctgact aatttttttt atttatgcag aggccgaggc cgcctcggcc tctgagctat     300
 110 tccagaagta gtgaggaggc ttttttggag gcctaggctt ttgcaaaaag ctcctcgagg     360
 111 aactgaaaaa ccagaaagtt aactggtaag tttagtcttt ttgtctttta tttcaggtcc     420
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 113 aggcctgtac ggaagtgtta cttctgctct aaaagctgct cgagtgtaaa acgacggcca     540
 114 gtacgtattt aatacgactc actatatggga atttaaatga attcggccgg ccgatatcct     600
 115 ggtaccgcgg ccgcggatct ccctttagtg agggttaatt ggatccagac atgataagat     660
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 119 cattaatgaa tcggccaacg cgcggggaga ggcggtttgc gtattgggcg ctcttccgct     900
 120 tcctcgctca ctgactcgct gcgctcggtc gttcggctgc ggcgagcggt atcagctcac     960
 121 tcaaaggcgg taatacggtt atccacagaa tcaggggata acgcaggaaa gaacatgtga    1020
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143 gtcagaagta agttggccgc agtgttatca ctcatggtta tggcagcact gcataattct 2340
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148 aactgatctt cagcatcttt tactttcacc agcgtttctg ggtgagcaaa aacaggaagg 2640
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157 agcccccgat ttagagcttg acggggaaag ccggcgaacg tggcgagaaa ggaagggaag 3180
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160 gaaacagcta tgaccatgat cctctagagt cgacctgcag gcatgcttaa ttaaggg 3357
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E--> 166 1/4

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 10/07/2005
PATENT APPLICATION: **US/10/550,788**   TIME: 09:31:07

Input Set : **A:\Sequence Listing 2005-1542A.txt**
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Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 3

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2

VERIFICATION SUMMARY   DATE: 10/07/2005
PATENT APPLICATION: **US/10/550,788**   TIME: 09:31:07

Input Set : **A:\Sequence Listing 2005-1542A.txt**
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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:166 M:254 E: No. of Bases conflict, LENGTH:Input:4 Counted:3358 SEQ:2
L:166 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:166 M:252 E: No. of Seq. differs, <211> LENGTH:Input:3357 Found:3358 SEQ:2